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AMENDMENTS TO THE SPECIFICATION

In the Specification:

Please replace the paragraph beginning on page 4, paragraph 44 with the following rewritten paragraph:

FIG 20: represents a sequence alignment of the membrane-proximal domains of vesiculovirus glycoproteins. The sequences shown are from the San Juan (SEQ ID No: 26) (37) and Orsay strains (SEQ ID No: 27) (19) of VSV Indiana, VSV New Jersey (SEQ ID No: 29) (18), Cocal virus (SEQ ID No: 28) (2), Chandipura virus (SEQ ID No: 31) (28), Piry virus (SEQ ID No: 30) (4) and spring viremia of carp virus (SVCV) (SEQ ID No: 32) (3). Residues in black colored font with light gray background are conserved among all the vesiculoviruses. Residues in white font with black background are identical residues in the virus sequences examined. Residues in black font with dark gray background indicate residues with similar properties. Stars at the bottom of the sequence represent invariant residues across the sequences examined.

In the Sequence Listing:

Please replace the sequence listing submitted on August 3, 2005 with the attached substitute sequence listing.